Network Systems
Science & Advanced
Computing

Biocomplexity Institute & Initiative

University of Virginia

Estimation of COVID-19 Impact in Virginia

July 15th, 2020

(data current to July 14th)

Biocomplexity Institute Technical report: TR 2020-088



BIOCOMPLEXITY INSTITUTE

biocomplexity.virginia.edu

Who We Are

- Biocomplexity Institute at the University of Virginia
 - Using big data and simulations to understand massively interactive systems and solve societal problems
- Over 20 years of crafting and analyzing infectious disease models
 - Pandemic response for Influenza, Ebola, Zika, and others



Points of Contact

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Biocomplexity COVID-19 Response Team

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Overview

• Goal: Understand impact of COVID-19 mitigations in Virginia

Approach:

- Calibrate explanatory mechanistic model to observed cases
- Project infections through the end of summer
- Consider a range of possible mitigation effects in "what-if" scenarios

Outcomes:

- Ill, Confirmed, Hospitalized, ICU, Ventilated, Death
- Geographic spread over time, case counts, healthcare burdens



Key Takeaways

Projecting future cases precisely is impossible and unnecessary. Even without perfect projections, we can confidently draw conclusions:

- Some VDH health districts have surging activity; which is pushing VA upward in the near term. Considering the experience of other states in nation, it is crucial to maintain control.
- Recent model updates:
 - Integrated "future Surge" scenarios as possible current scenarios
 - Identification and timing of districts experiencing a "Surge" developed
 - "Best fitting" scenarios per health district now include surging districts
 - Updated additional analyses to act as early indicators of surge and provide evidence for those surging
- Much of nation shows rapid rise following relaxation of social distancing with limited control measures.
- The situation is changing rapidly. Models will be updated regularly.

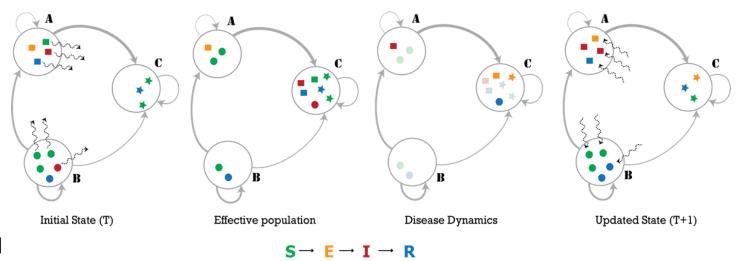


Model Configuration and Data Analysis



Simulation Engine – PatchSim

- Metapopulation model
 - Represents each population and its interactions as a single patch
 - 133 patches for Virginia counties and independent cities
- Extended SEIR disease representation
 - Includes asymptomatic infections and treatments
- Mitigations affect both disease dynamics and population interactions
- Runs fast on high-performance computers
 - Ideal for calibration and optimization





Susceptible → Exposed → Infectious → Removed

Venkatramanan, Srinivasan, et al. "Optimizing spatial allocation of seasonal influenza vaccine under temporal constraints." PLoS Computational Biology 15.9 (2019): e1007111.

Model Configuration

- Transmission: Parameters are calibrated to the observed case counts
 - Reproductive number: 2.1 2.3
 - Infectious period (time of infectiousness before full isolation): 3.3 to 5 days
- Initial infections: Start infections from confirmed cases by county
 - Timing and location based on onset of illness from VDH data
 - Assume 15% detection rate, so one confirmed case becomes ~7 initial infections
- Mitigations: Intensity of social distancing rebound and control sustaining mitigations into the future are unknowable, thus explored through 5 scenarios



Full Model Parameters

	Parameter	Values	Description
Transmission	Transmissibility (R ₀) ¹	2.2 [2.1 – 2.3]	Reproductive number
	Incubation period ¹	5 days	Time from infection to infectious
	Infectious period ¹	3.3 - 5 days	Duration of infectiousness
Tran	Infection detection rate ³	15%	1 confirmed case becomes ~7 initial infections
_	Percent asymptomatic ¹	50%	Infected individuals that don't exhibit symptoms
Resources	Onset to hospitalization ¹	5 days	Time from symptoms to hospitalization
	Hospitalization to ventilation ¹	3 days	Time from hospitalization to ventilation
	Duration hospitalized	8 days	Time spent in the hospital ⁴
	Duration ventilated ²	14 days	Time spent on a ventilator
	Percent hospitalized ¹	5.5% (~20% of confirmed)	Symptomatic individuals becoming hospitalized
	Percent in ICU ¹	20%	Hospitalized patients that require ICU
	Percent ventilated ¹	70%	ICU patients requiring ventilation
	Percent Fatality	1.35%	Symptomatic individuals who die

¹ CDC COVID-19 Modeling Team. "Best Guess" scenario. Planning Parameters for COVID-19 Outbreak Scenarios. Version: 2020-03-31.

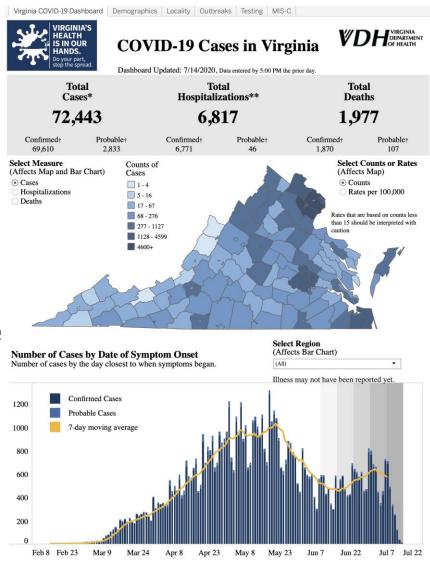
² Up-to-date. COVID-19 Critical Care Issues. https://www.uptodate.com/contents/coronavirus-disease-2019-covid-19-critical-care-issues?source=related_link (Accessed 13APRIL2020)

³ Li et al., Science 16 Mar 2020:eabb3221 https://science.sciencemag.org/content/early/2020/03/24/science.abb3221 (Accessed 13APRIL2020)

⁴ Personal communications, UVA Health and Sentara ($^{\sim}$ 500 VA based COVID patients) 16-Jul-20

Calibration Approach

- Data:
 - County level case counts by date of onset (from VDH)
 - Confirmed cases for model fitting
- Model: PatchSim initialized with disease parameter ranges from literature
- Calibration: fit model to observed data
 - Search transmissibility and duration of infectiousness
 - Markov Chain Monte Carlo (MCMC) particle filtering finds best fits while capturing uncertainty in parameter estimates
- **Spatial Adjustments:** VDH districts grouped to 3 tiers of growth during the Pause, with similarly scaled reductions then applied to the groups of districts
- **Project:** future cases and outcomes using the trained particles



Accessed 2pm July 14, 2020 https://www.vdh.virginia.gov/coronavirus/

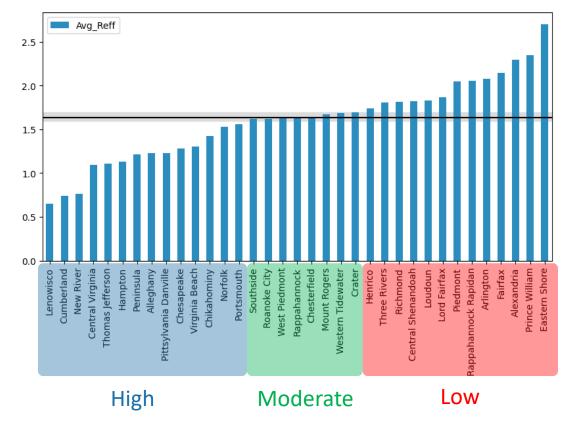
Spatial Adjustments at District Level

District Specific adjustments based on Growth

during Pause

• Group districts by their mean growth from mid-April to mid-May (using model based $R_{\rm eff}$)

- Assign reductions during Pause, and beyond, to members of these groups
- Low reduction = 40%
- Moderate reduction = 45% (previous level)
- High reduction = 55%





Scenarios: Past to Present

Pause from Social Distancing: Began on March 15th

- Lifted on May 15th (61 days), with two-week delay (75 days) for select counties*
- **Intensity**: Social distancing pauses and <u>significantly reduces case growth</u>, this level varies by VDH Health District and is fit through an analysis of growth rate during the Pause

Intensity of Rebound: Some districts rebounded following initial relaxation of Pause

- Steady: Intensity of effective mixing remains steady from Pause as infection control practices moderate increased interactions
- **Light:** Effective mixing returns to 1/6th of pre-pandemic levels
- Full Rebound: Interactions return completely (100%) to pre-pandemic levels, as a reference

Tracing and Isolation: Increased Testing Capacity coupled with infection control measures can limit the period of infectiousness without isolation

• **Better Detection**: Observed relative reductions in days from onset to diagnosis applied to infectious period from $(30\% \rightarrow 45\% \rightarrow 30\%)$ and remain stable into future for projections



Mitigation Scenarios: Present to Future

Resurgence: Much of the nation experiencing a resurgence

- Some districts in the Commonwealth also showing a resurgence
- 23 states surging: 28-day delay (avg) from relaxation to surge

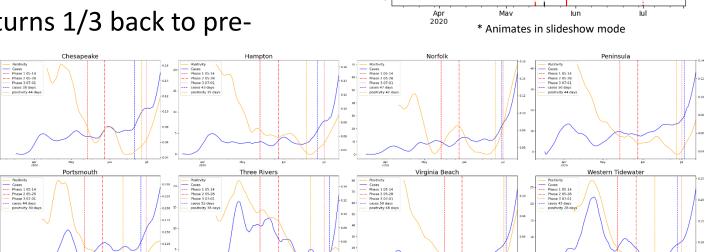
Intensity of Surge: Difficult to predict with limited data

Strong Rebound: Effective mixing returns 1/3 back to pre-

pandemic levels

Timing of Surge: Present and Future

- Determine surging districts and timing through "hockey stick" fit
- Allow "Best Fit" method to select from "Surge" scenarios
- Default to 28 days from July 1st for districts without present surge



RSS: 609.072

Portsmouth

Eight districts in Eastern region with surging incidence (mid-June to early July)

** Update latest run Wed AM, Thomas Jefferson & Pittsylvania-Danville tip over to surge



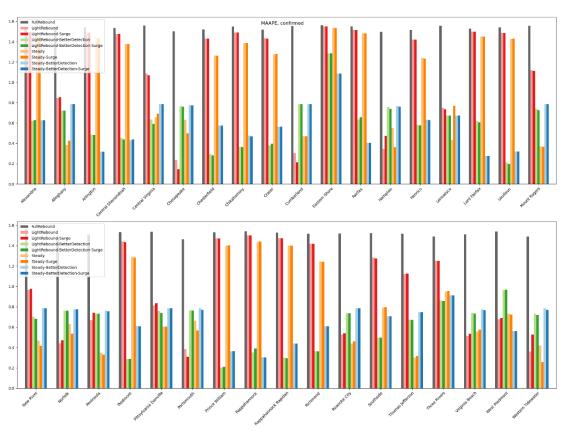
Eight Scenarios for Projection

Abbr	Rebound Intensity	Better Detection	Surge	Name
LR	Light	No	No	LightRebound
LR-S	Light	No	Yes	LightRebound-Surge
LR-BD	Light	Yes	No	LightRebound-BetterDetection
LR-BD-S	Light	Yes	Yes	LightRebound-BetterDetection-Surge
S	Steady	No	No	Steady
S-S	Steady	No	Yes	Steady-Surge
S-BD	Steady	Yes	No	Steady-BetterDetection
S-BD-S	Steady	Yes	Yes	Steady-BetterDetection-Surge



Selection of Best Fitting Projection

Recent incidence by district (last week) is measured against all eight projections, one with least error is selected as the "Best Fit" projection



Abbr	Name	# of Districts (last wk)
LR	LightRebound	2 (2)
LR-S	LightRebound-Surge	3 (0)
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S	Steady	6 (12)
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S-BD	Steady-BetterDetection	5 (7)

- 10 districts have Surge projections as BestFit
- Continued movement towards "higher incidence" projections



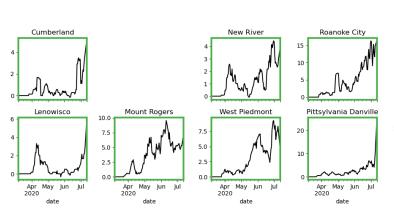
Data Analysis Supporting Model

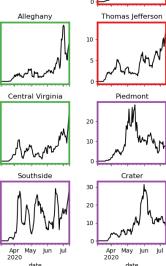


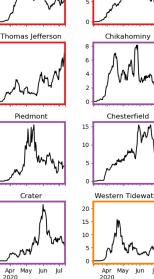
Case Rate (per 100k) by VDH District

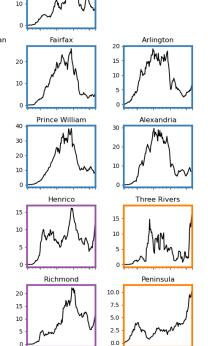
Sharp increases in some health districts

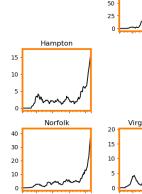
- Regions arranged to rough position in Commonwealth and colored by VDH Health Region
- Considerable variation across districts
- Some consistent behaviors during mid-April to mid-May during the Pause period
- Smoothed (Savitzky-Golay filter)















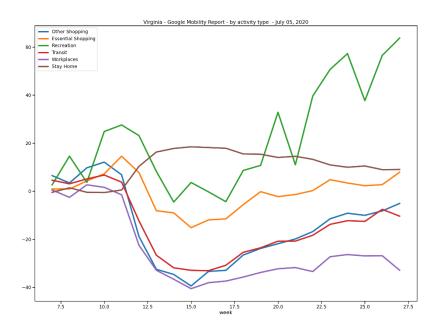
Estimating Effects of Social Distancing

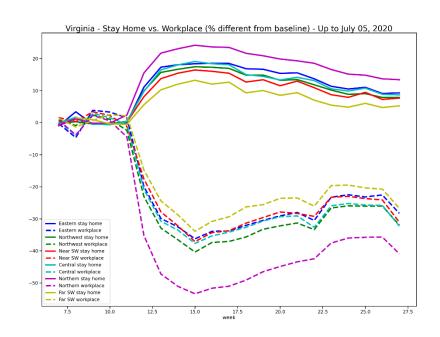
Mobility data shows pause mid-March, slow rebound starting in May

Google Mobility data shows continued slow rebound (as of July 5th)

https://www.google.com/covid19/mobility/

- Continued reduction of those staying at home, very slow and stable reductions
- Other activities show vaster increases with grocery / retail nearly back to baseline
- Parks and recreation show significant increase





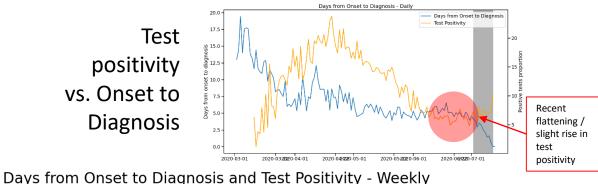
Changes in Case Detection

VDH data show changes in time from Symptom Onset to Diagnosis

Days to Diagnosis dropped but rebounding

- Mid March to Late April = 7.8 days
- Late April to Mid May = 5.8 days (25% lower)
- Mid May to early June = 4.8 days (39% lower)

• Early June to late June = 5.2 days (33% lower)



Days from Onset to Diagnosis Weekly PCR Test Positivity

17.5

15.0 4 5 10.0 Days from Onset to Diagnosis Weekly PCR Test Positivity

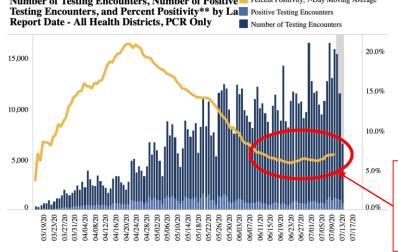
17.5

15.0 4 5 10.0 Days from Onset to Diagnosis Weekly PCR Test Positivity

17.5

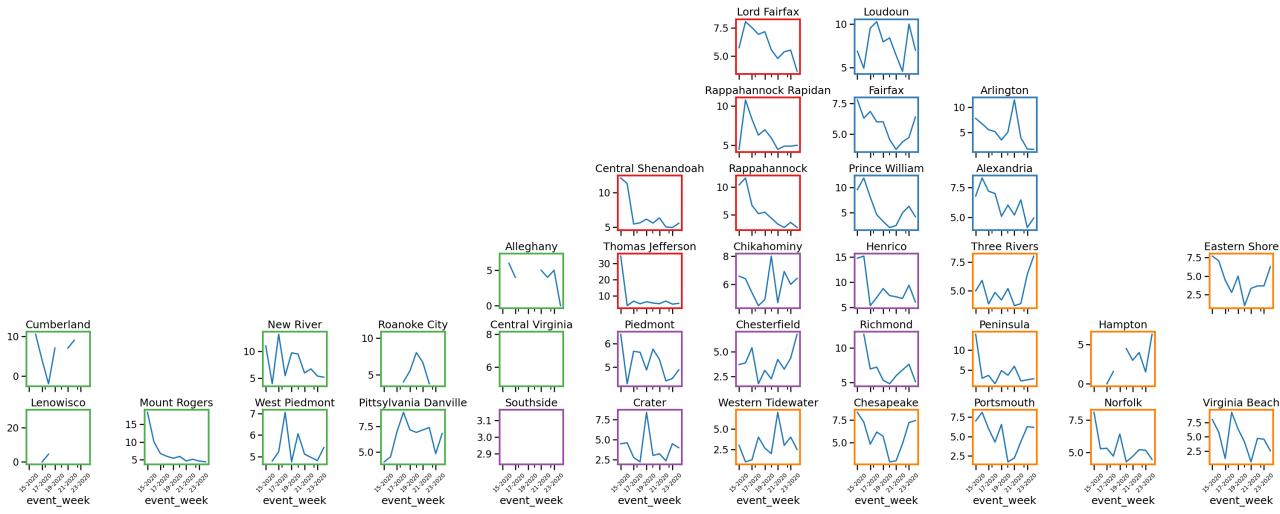
15.0 4 5 10.0 Days from Onset to Diagnosis Days from Onset Days from On





Recent flattening then slight rise in total tests

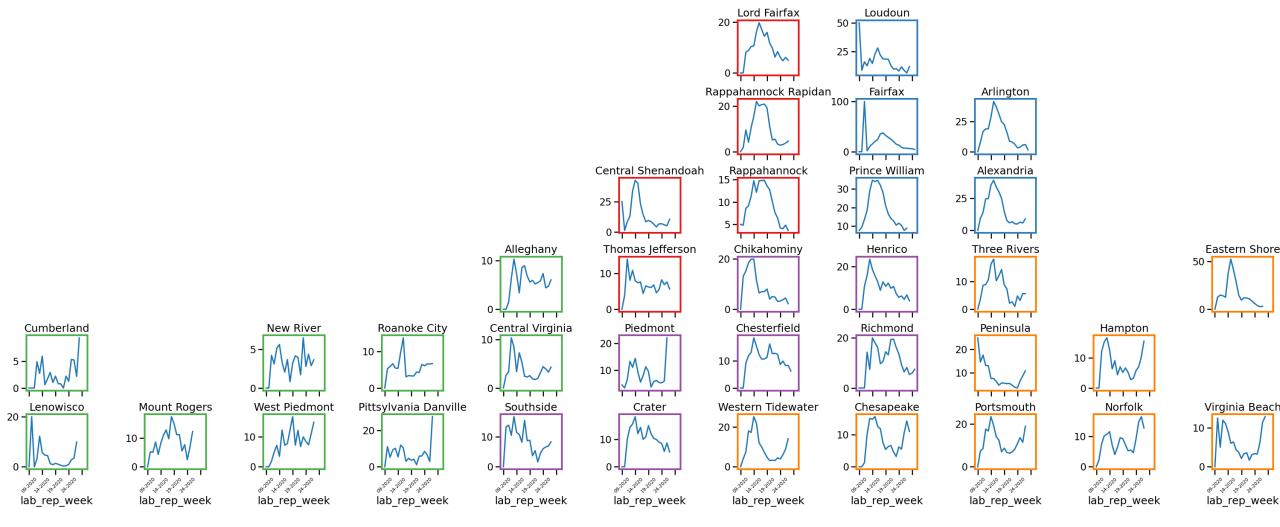
Changes in Case Detection* – by district



^{*}up to the early July when data is stable

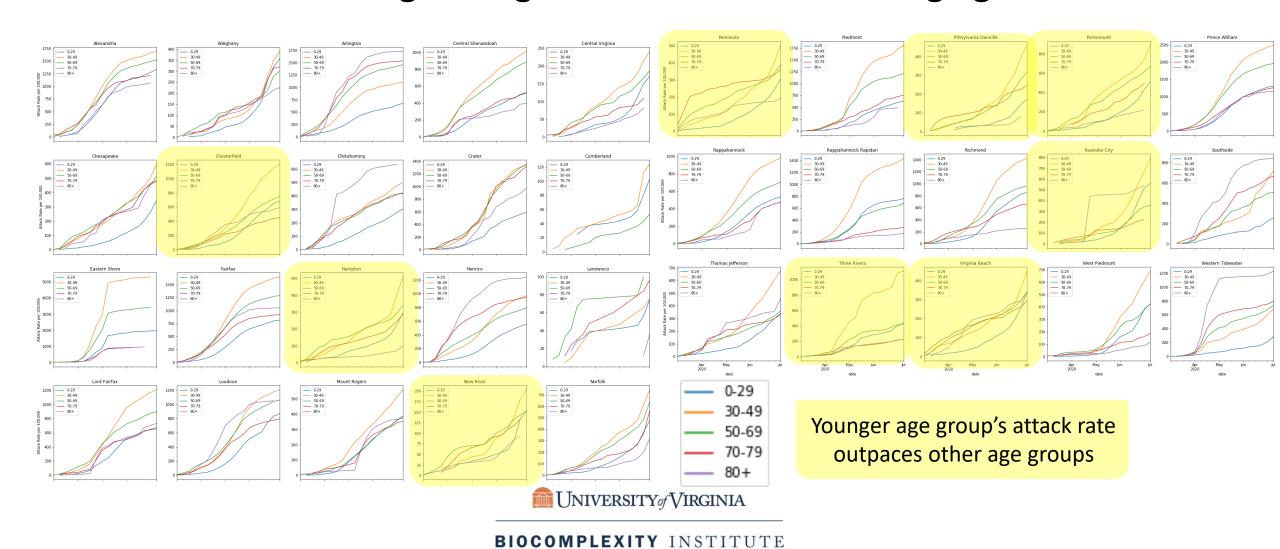


Changes in Test Positivity – by district



Case Attack rate (per 100k) by Age-group and District

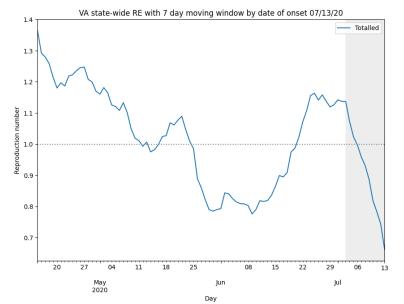
VDH data show changes in age attack rates in some surging districts

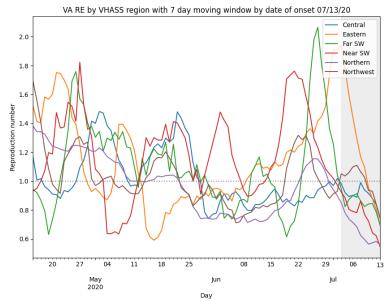


Estimating Daily Reproductive Number

July 3rd Estimates

Region	Current R _e	Diff Last Week
State-wide	1.137	0.013
Central	0.983	0.072
Eastern	1.940	0.540
Far SW	0.949	-0.986
Near SW	0.924	-0.305
Northern	0.852	-0.252
Northwest	1.050	0.057



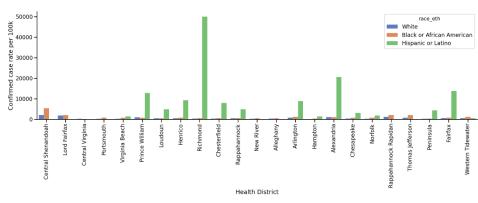


Methodology

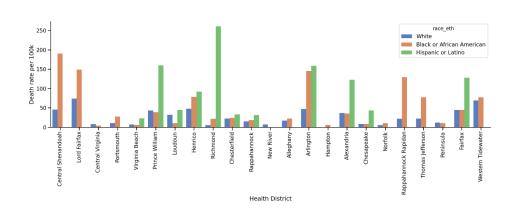
- Wallinga-Teunis method as implemented in EpiEstim¹ R package
- Based on Date of Onset of Symptoms
- Uses serial interval to estimate R_e over time: 6 days (2 day std dev)
 Recent Estimates subject to revision as more data comes in
- Date of onset unstable in last 7-14 days

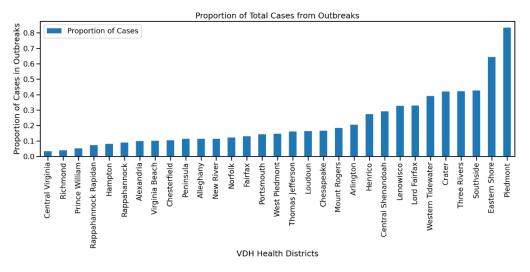
Impact of Race / Ethnicity & Outbreaks

Confirmed Case Rate



Death Rate





Different Races and Ethnicities disproportionally affected

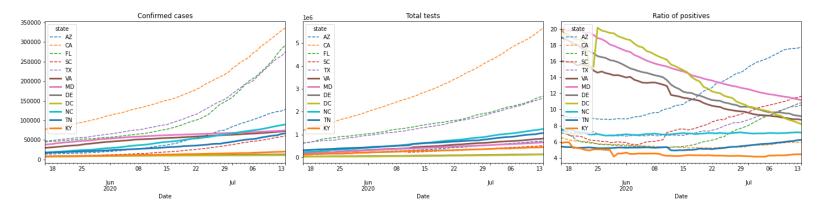
 Hispanic population bears large burden of disease compared to population size

Outbreak Events are hard to predict and affect model fits

- Eastern Shore has 60% of cases from 10 outbreaks
- Fairfax most outbreaks but relatively low proportion



Other State Comparisons



Several States continue to experience large surges in cases

 Some of these surging states show no signs of abating, however, several out west may be slowing their case incidence

R_e Estimates for VA and neighbors show upward trend

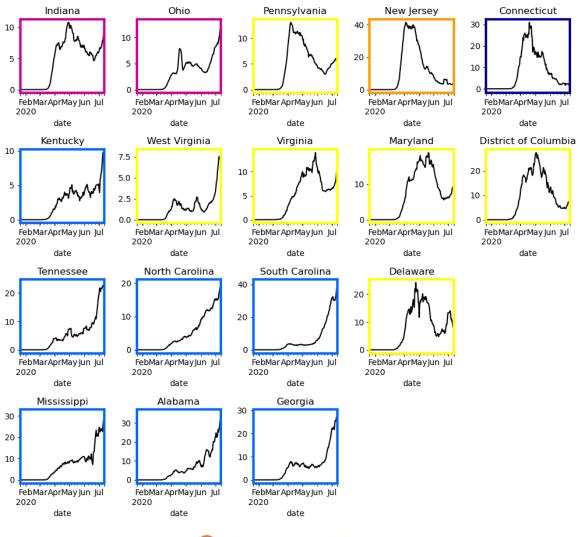
- Virginia above 1, as are all but DC
- Tennessee and Delaware experiencing sharp rise in R

Signs of resurgence: Plateauing or increase in test positivity and R above 1 for several weeks

Estimated R_e* for surging States and Neighbors



Other State Comparisons

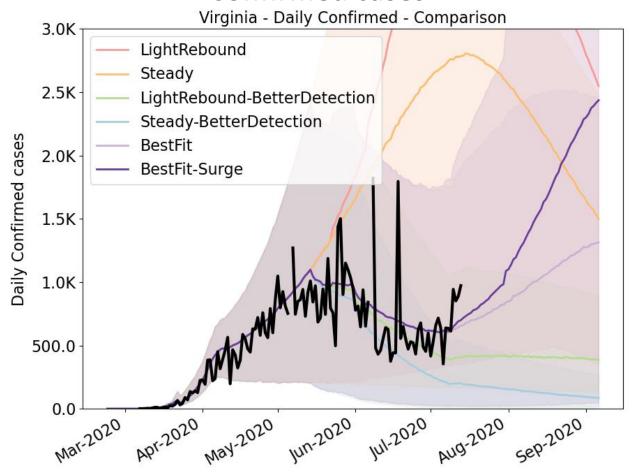


Model Results

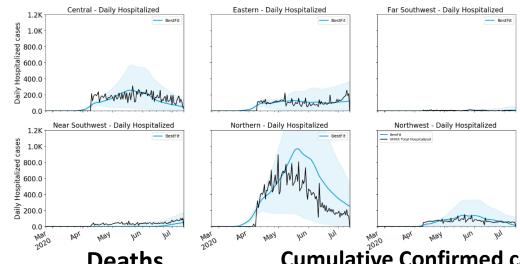


Outcome Projections

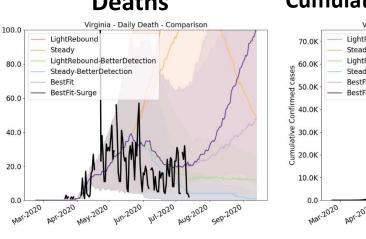
Confirmed cases



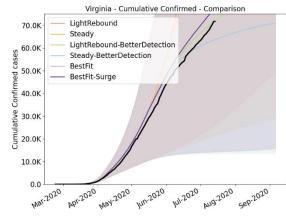
Hospital occupancy



Deaths

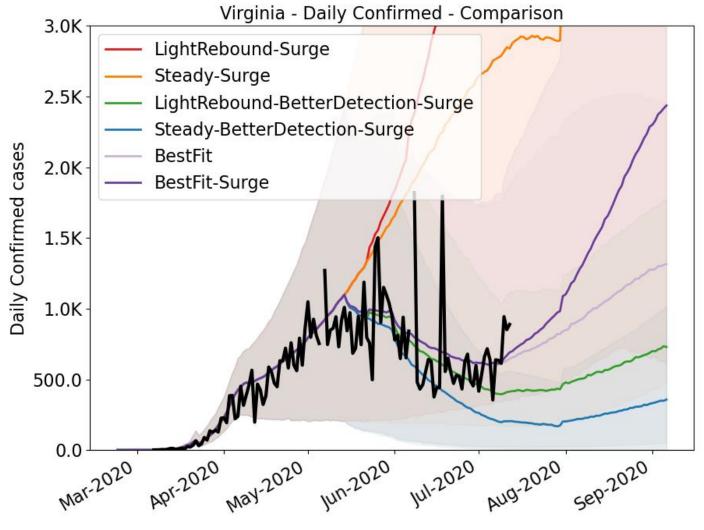


Cumulative Confirmed cases





Outcome Projections – with Surge



Weekly New Confirmed Cases*

Week Ending	Best Fit	Best Fit w/ Surge
7/5/20	4,311	4,330
7/12/20	4,254	4,316
7/19/20	4,650	5,010
7/26/20	5,076	5,734
8/2/20	5,612	6,728
8/9/20	6,239	8,440
8/16/20	6,958	10,267
8/23/20	7,639	12,220
8/30/20	8,340	14,320
9/6/20	8,926	16,121

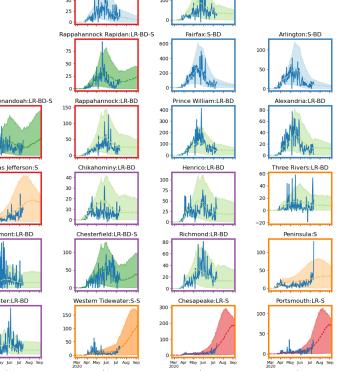


^{*}Numbers are medians of projections

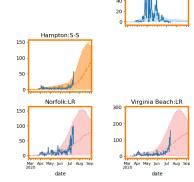
District Level Projections - Daily

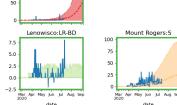
Best fitting projections by District

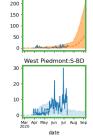
- Projections that best fit recent trends
- Daily confirmed cases by Region (blue solid) with simulation at the region level (black dotted)
- Projection color consistent with other and abbreviated in title

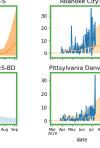


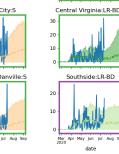
Abbr	Name	# of Districts (last wk)
LR	LightRebound	2 (2)
LR-S	LightRebound-Surge	3 (0)
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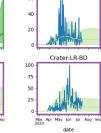


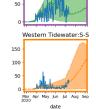




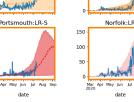












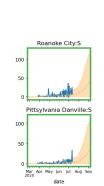


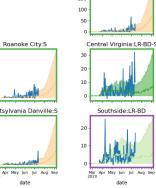


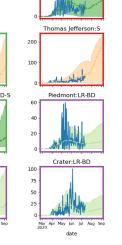
District Level Projections – Daily with Surge

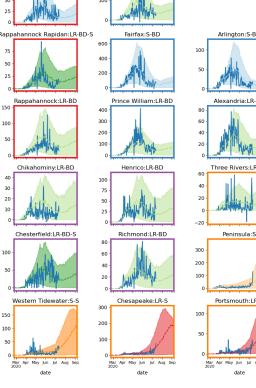
Best fitting projections by District

- Projections that best fit recent trends with Surge assumed for all districts
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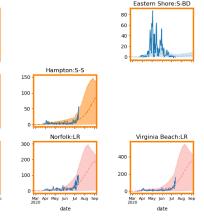








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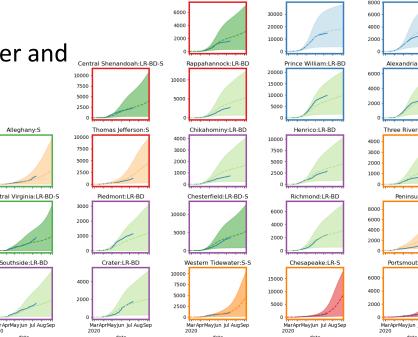




District Level Projections - Cumulative

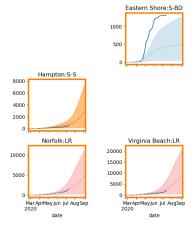
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10000

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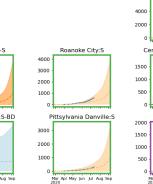


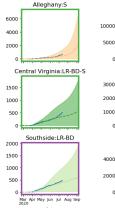
Cumberland:LR-

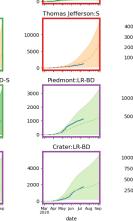
District Level Projections – Cumulative with Surge

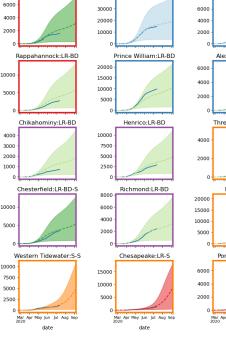
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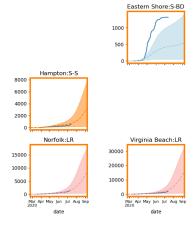






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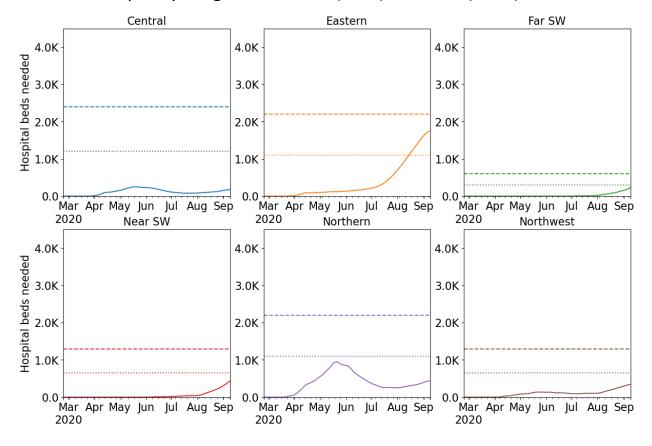




Hospital Demand and Capacity by Region

Capacities by Region – BestFit- Surge

COVID-19 capacity ranges from 80% (dots) to 120% (dash) of total beds



- Based on current best fits with potential surge
 - Eastern region exceeds 80% capacity in mid-August
 - Multiple regions (Near SW, Far SW, Eastern and Northern) may near their capacity in September
- Next few weeks (until mid-August) are crucial to mitigate/prepare for a surge in cases
- Activity in neighboring states and reopening of schools/universities may make this more likely

* Assumes average length of stay of 8 days

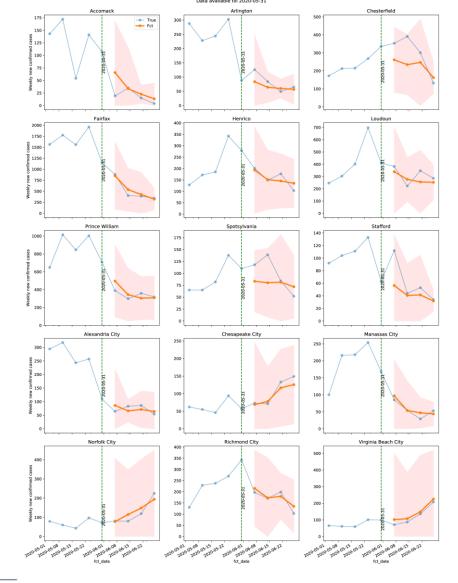


A new forecasting framework nearing completion

- Augmenting "forecast through projection selection" with ensemble of statistical and mechanistic models
- Methods included:
 - Autoregressive methods with mobility data, google trends, other county case counts, and weather data as exogenous regressors.
 - Long short-term memory deep learning models with additional mobility data, Google search trends, weather data for training.
 - Mechanistic models for relative ease of incorporation of intervention scenarios.
- Forecasts from multiple models (methods) combined to yield probabilistic forecasts: Bayesian Model Averaging

$$P(y|f_1, f_2, ..., f_M) = \sum_{m=1}^{M} w_m g_m(y|f_m)$$

 $y = \text{true value}, f_m = \text{forecast from } m^{th} \text{ model}$ $g_m(y|f_m) = m^{th} \text{ model probabilistic forecast}$ $w_m = \text{weights assigned to } m^{th} \text{ model forecast}$



Key Takeaways

Projecting future cases precisely is impossible and unnecessary. Even without perfect projections, we can confidently draw conclusions:

- Some VDH health districts have surging activity; which is pushing VA upward in the near term.
 Considering the experience of other states in nation, it is crucial to maintain control.
- Recent model updates:
 - Integrated "future Surge" scenarios as possible current scenarios
 - Identification and timing of districts experiencing a "Surge" developed
 - "Best fitting" scenarios per health district now include surging districts
 - Updated additional analyses to act as early indicators of surge and provide evidence for those surging
- Much of nation shows rapid rise following relaxation of social distancing with limited control measures.
- The situation is changing rapidly. Models will be updated regularly.



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Biocomplexity Institute. COVID-19 Surveillance Dashboard. https://nssac.bii.virginia.edu/covid-19/dashboard/

Google. COVID-19 community mobility reports. https://www.google.com/covid19/mobility/

Cuebiq: COVID-19 Mobility insights. https://www.cuebiq.com/visitation-insights-covid19/

Biocomplexity page for data and other resources related to COVID-19: https://covid19.biocomplexity.virginia.edu/



Questions?

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Supplemental Slides



Recent Parameter Validation

New York State announced sero-prevalence survey results on May 2nd

- 15,000 antibody tests conducted randomly through the state at grocery stores
- Total Attack Rate: 12.3%

Estimation of undetected infections

- Total infections in NY = 2.46M, total of 300K confirmed cases
- Confirmed case detection = 12% of infections (close to 15% used in model)

Estimation of hospitalizations from infections

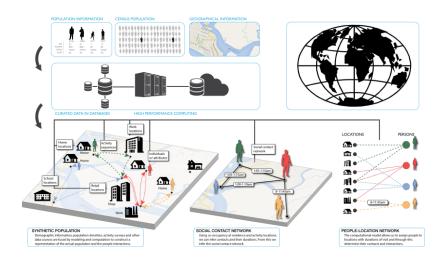
- Total infections in NY = 2.46M, total of 66K hospitalizations
- Hospitalizations = 2.7% of infections (close to 2.25% used in model)



Agent-based Model (ABM)

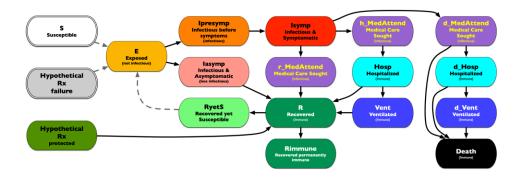
EpiHiper: Distributed network-based stochastic disease transmission simulations

- Assess the impact on transmission under different conditions
- Assess the impacts of contact tracing



Synthetic Population

- Census derived age and household structure
- Time-Use survey driven activities at appropriate locations



Detailed Disease Course of COVID-19

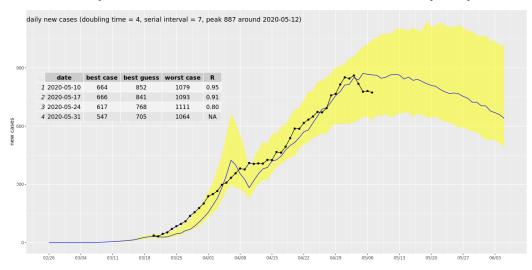
- Literature based probabilities of outcomes with appropriate delays
- Varying levels of infectiousness
- Hypothetical treatments for future developments



ABM Social Distancing Rebound Study Design

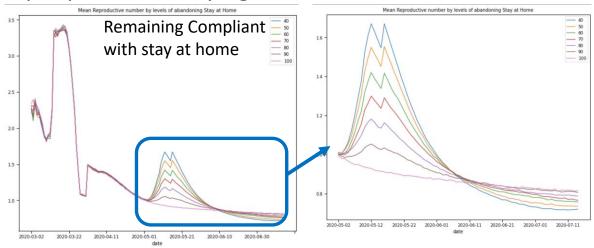
Study of "Stay Home" policy adherence

- Calibration to current state in epidemic
- Implement "release" of different proportions of people from "staying at home"



Calibration to Current State

- Adjust transmission and adherence to current policies to current observations
- For Virginia, with same seeding approach as PatchSim



Impacts on Reproductive number with release

- After release, spike in transmission driven by additional interactions at work, retail, and other
- At 25% release (70-80% remain compliant)
- Translates to 15% increase in transmission, which represents a 1/6th return to pre-pandemic levels



Medical Resource Demand Dashboard

https://nssac.bii.virginia.edu/covid-19/vmrddash/

